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Deletion of the mouse *RegIII β* (Reg2) gene disrupts ciliary neurotrophic factor signaling and delays myelination of mouse cranial motor neurons

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A large number of cytokines and growth factors support the development and subsequent maintenance of postnatal motor neurons. *RegIII β* , also known as *Reg2* in rat and *HIP/PAP1* in humans, is a member of a family of growth factors found in many areas of the body and previously shown to play an important role in both the development and regeneration of subsets of motor neurons. It has been suggested that *RegIII β* expressed by motor neurons is both an obligatory intermediate in the downstream signaling of the leukemia inhibitory factor/ciliary neurotrophic factor (CNTF) family of cytokines, maintaining the integrity of motor neurons during development, as well as a powerful influence on Schwann cell growth during regeneration of the peripheral nerve. Here we report that in mice with a deletion of the *RegIII β* gene, motor neuron survival was unaffected up to 28 weeks after birth. However, there was no CNTF-mediated rescue of neonatal facial motor neurons after axotomy in KO animals when compared with wild-type. In mice, *RegIII β* positive motor neurons are concentrated in cranial motor nuclei that are involved in the patterning of swallowing and suckling. We found that suckling was impaired in *RegIII β* KO mice and correlated this with a significant delay in myelination of the hypoglossal nerve. In summary, we propose that *RegIII β* has an important role to play in the developmental fine-tuning of neonatal motor behaviors mediating the response to peripherally derived cytokines and growth factors and regulating the myelination of motor axons.

Schwann cells | suckling | hypoglossal nerve | LIF

A large number of neurotrophic factors and cytokines have been shown to sustain developing motor neurons and to encourage the survival of postnatal motor neurons after damage (1–6). Rat *Reg2* [also known as *RegIII β* in mouse and *HIP/PAP1* in humans (7)] has been reported to play an important role in both the support of motor neurons during development and in the process of axon regeneration through axon–Schwann cell signaling in the adult peripheral nervous system (8, 9). *Reg2* is a member of a large family of over 17 related genes divided into four subtypes (types I, II, III, and IV) based on the primary structures of the encoded proteins of the genes (10–13). *Reg2* is the equivalent of mouse *RegIII β* gene, which share 90% homology at both the nucleotide and protein level. First identified as a transcript up-regulated in pancreatitis, *Reg2*, a secreted protein (relative *M_r* 16,000) found in many sites throughout the body, was shown to have an anti-apoptotic action on pancreatic cell lines (14). *Reg2* also promotes the growth of epithelial intestinal cells, whereas loss of *Reg2/RegIII β* delayed liver regeneration (10, 15–18).

Two roles have been proposed for *Reg2* in the nervous system. First, *in vivo* studies have suggested that *Reg2* is released from

damaged motor and sensory neurons and has a proregenerative function (9, 19, 20). Secondly, *in vitro* data have cast *Reg2* as a neurotrophic factor for motor neurons acting as an obligatory intermediate for the ciliary neurotrophic factor (CNTF)/leukemia inhibitory factor (LIF) family of cytokines (8). In the adult rat, *Reg2* is massively up-regulated in motor neurons and some sensory neurons after nerve crush (19) and is rapidly transported to the lesion site, where it is thought to be secreted and act on Schwann cells. *In vitro*, *Reg2* has a mitogenic effect on Schwann cells and inhibition of *Reg2* with a neutralizing antibody retards the progress of regeneration. Taken together, it seemed likely that the Schwann cell response at the point of axotomy was potentiated by release of *Reg2* from damaged axons (9).

Here, we describe the effects of targeted ablation of the *RegIII β* gene in mice and show a developmental role of *RegIII β* in axon–Schwann cell signaling. We also report that *RegIII β* is required to promote the response of subsets of motor neurons to CNTF but not for motor neuron survival.

Results

Generation of *RegIII β* -Deficient Mice. KO mice were homozygous for a targeted disruption of the *RegIII β* gene locus created in ES cells, in which a region from exons 2 to 5 was deleted and replaced by an IRES-*Tau-LacZ-loxP/MC1neopA/loxP* reporter/selection cassette, thus resulting in a null allele. Mice homozygous for the *RegIII β* -null allele were phenotypically indistinguishable from wild-type or heterozygous littermates. No embryonic lethality or significant developmental defects were observed in *RegIII β* ^{−/−} animals. Adult *RegIII β* ^{−/−} mice of both sexes were fertile, and litter size was normal. We compared the expression of *RegIII β* between wild-type and KO animals by using quantitative RT-PCR (RT-qPCR) and found that, as expected, the expression of *RegIII β* mRNA dropped from 100 ± 25.2% (wild type) to 0.08 ± 0.02% (KO) (*n* = 6 per group). Immunohistochemistry also showed a complete lack of *RegIII β* protein-like immunoreactivity in postnatal KO mice (Fig. 1). To check for potential compensatory up-regulation, we also mea-

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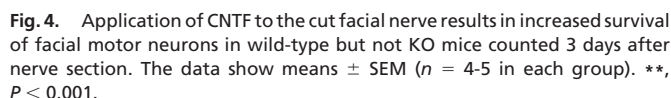
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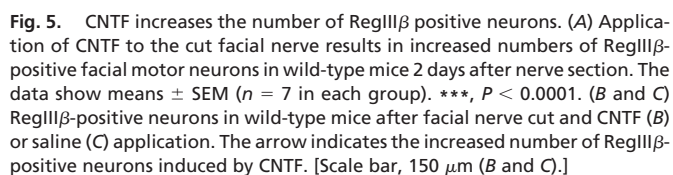
We have previously shown (9) that section of the sciatic nerve results in down-regulation of Reg2/RegIII β in motor neurons in rat pups and up-regulation in adult rats. However, in mouse pups with facial nerve axotomy at P3.5 and perfused 2 days later, we did not detect any change in numbers of RegIII β -positive neurons in the facial motor nucleus. However, application of CNTF after facial nerve section at P3.5 in mouse pups significantly increased the number of RegIII β -positive neurons seen 2 days later to 148% compared with the contralateral side ($P < 0.0001$) (Fig. 5 and Fig. S3). Finally, whereas in adult rats, Reg2 is expressed by all facial motor neurons within 24h of axotomy, in adult mice, RegIII β is not reexpressed in facial motor neurons 1–7 days after axotomy (Fig. S4).

to be concentrated in cranial motor neurons that have been associated with suckling and swallowing, such as the hypoglossal nucleus and nucleus ambiguus. We therefore examined the ability of RegIII β KO mice to ingest milk during a 1-h period of suckling after a 2-h period of isolation from the mother. Compared with wild-type pups, we found a significant reduction in milk/colostrum ingestion: percentage weight increase in wild-type mice, $3.1 \pm 0.2\%$; and in KO mice, 0.45 ± 0.2 ($P < 0.01$).

The impaired suckling phenotype, thus, may result from a disruption of myelination in mutant mice. We also examined the hypoglossal innervation of tongue musculature in wild-type and KO mice by using protein gene product (PGP) as a marker for nerve fibers and α -bungarotoxin for muscle end plates. We found no evidence of changes in innervation density or size of muscle end plates (data not shown).

Previous research has suggested that *in vitro* rat Reg2 is a motor neuron survival factor essential for the actions of CNTF-like cytokines and a powerful Schwann cell mitogen that potentiates axonal repair and regeneration *in vivo* (8, 9). However, we show here that in mice with a genetic deletion of the RegIII β gene (the equivalent gene to Reg2 in rats), there is no increased motor neuron cell death during development. Nevertheless, we report that the efficacy of CNTF in reducing neuronal cell death was diminished in RegIII β KO mice indicating that RegIII β is important for the survival functions of CNTF. Finally, we found that myelination was delayed in subsets of hypoglossal motor neurones in KO animals and the efficiency of milk ingestion reduced.

RegIII β Is an Intermediate in the CNTF Survival Pathway. The suggestion that Reg2/RegIII β was a motor neuron neurotrophic factor and a signaling intermediary in the CNTF survival pathway stems from elegant *in vitro* studies demonstrating that purified Reg2 can act as a paracrine/autocrine neurotrophic factor for a subset of motor neurons (8). Furthermore, it has been shown that CNTF, as well as the related factors LIF, cardiotrophin 1, and oncostatin-M, rapidly induces Reg2 mRNA in some motor neurons and that released Reg2 acts in a paracrine/autocrine fashion to support motor neurons (8, 9). The CNTF/LIF family of cytokines signal through a receptor complex that includes the LIF receptor (LIFR) subunit, and it has been



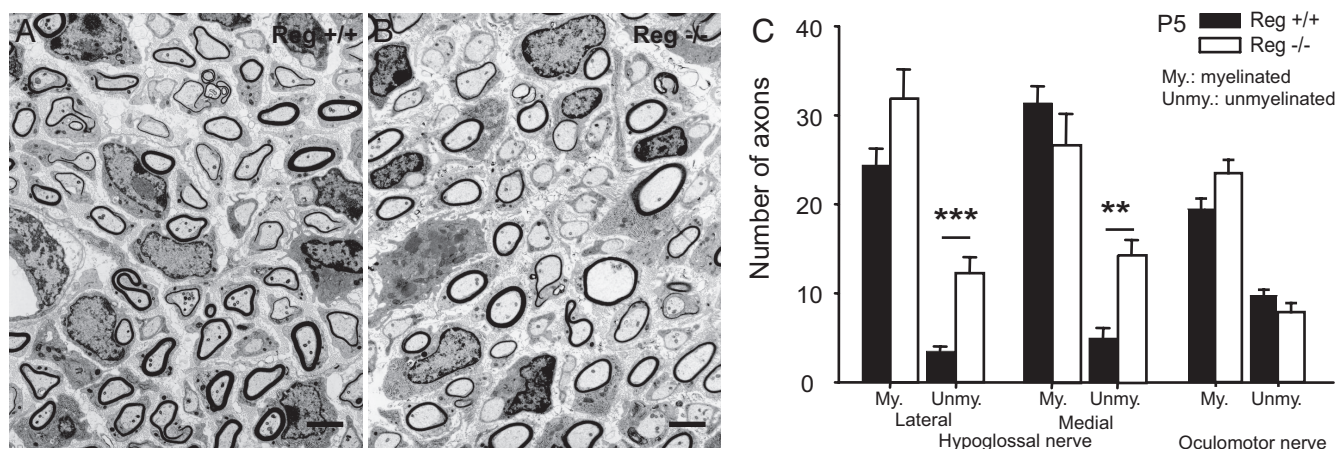


Fig. 6. Increased number of unmyelinated axons in RegIII β KO mice. Electron micrograph of the medial hypoglossal nerve of RegIII β wild-type (A) or RegIII β KO (B) mice. There is an increase in the number of unmyelinated fibers in the lateral and medial hypoglossal motor nerve of RegIII β KO mice ($n = 8$ in each group). (C) No differences were observed in the oculomotor nerve ($n = 3$ in each group). The data show means \pm SEM. **, $P < 0.001$; ***, $P < 0.0001$. All nerves were studied at P5. [Scale bar, 2 μ m (A and B).]

shown previously that in LIFR KO mice, there is no expression of Reg2/RegIII β during development (9). Thus, Reg2/RegIII β expression in some developing motor (and sensory) neurons depends on cytokines of the LIF family acting through a receptor complex containing LIFR.

The absence of motor neuron cell death in KO mice, therefore, is puzzling, especially given that *RegIII β* is essential for the *in vitro* neuronal survival effects of CNTF to be manifest. The evidence for a lack of effect on motor neuron survival comes from the presence of β -galactosidase reaction product in postnatal neurons up to P10 and the normal numbers of facial motor neurons in KO mice. However, in CNTF KO mice, there is similarly no increased loss of motor neurons during the first weeks of life (24, 25). This, perhaps, is not surprising, given that levels of CNTF are low during development and, although Schwann cells are the richest source of CNTF in the adult peripheral nervous system, levels only rise during the first postnatal week (26–28). Knockout of the CNTF gene did result in motor neuron loss 28 weeks after birth (24); however, *RegIII β* mice did not show motor neuron loss in later adult stages.

Previous studies suggest that CNTF itself is not the key ligand acting at the CNTF/LIF receptor complex. Nishimune *et al.* (8) found that RegIII β expression was unimpaired or delayed in a range of KOs including *cntf*, *ct1*, and *cntf/lif* double mutants, leading them to suggest that a factor as yet unknown was key to driving the expression of RegIII β in motor neurons.

That CNTF-like factors are as important *in vivo* as *in vitro* and require $\text{RegIII}\beta$ expression is implied by the observation that $\text{RegIII}\beta$ expression increases in axotomized neurons only when CNTF is applied to the nerve stump and that CNTF has no survival effect on motor neurons in $\text{RegIII}\beta$ KO mice. We also show that the number of $\text{RegIII}\beta$ neurons increased in wild-type mice when CNTF was applied to the cut nerve. This implies that many motor neurons have the capacity to express $\text{RegIII}\beta$, but this number is restricted by availability of CNTF-like factors during development. Nevertheless, $\text{RegIII}\beta$ may not be the only intermediary involved in CNTF-like factor signaling as only $\approx 15\text{--}20\%$ of facial motor neurons express $\text{RegIII}\beta$, and CNTF has previously been shown to rescue $\approx 75\%$ of rat facial motor neurons from cell death when axotomized soon after birth (24). However, the substantially larger concentration of CNTF used in these experiments ($5\text{ }\mu\text{g}$ vs. 250 ng used in the present study) may well have driven the expression of $\text{RegIII}\beta$ in many more axotomized facial motor neurons than seen here and resulted in greater levels of survival (24). Also the dynamics of Reg2

expression in rat are different from that of RegIII β in mice. RegIII β never reappeared in the adult mouse after section of the facial nerve (Fig. S4) or sciatic nerve (data not shown) and did not noticeably decrease 24 h after axotomy, as was observed after sciatic transection in neonatal rats. Presumably other factors may play a similar role to RegIII β in adult mouse motor neurons, although it is unlikely to be RegIII α , which although up-regulated in the KO mouse, did not potentiate CNTF-mediated survival.

Growth Factors and Myelination. We show here that myelination of a subset of hypoglossal motor neuron axons is delayed in RegIII β KO mice at P5.5 but normal by P21. This seems likely to be attributable to the loss of RegIII β because myelination of oculomotor axons in KO mice was normal and oculomotor motor neurons did not express RegIII β at any stages of development. This delayed myelination would be expected to disrupt the transmission of signals along the axon and may account for the reduced efficiency of suckling behavior. Delayed myelination did not, however, result in any reduction in the size or number of motor neuron end plates in target muscles of the hypoglossal such as the glossohyoid (unpublished data). However, how then does RegIII β promote myelination in discrete populations of motor neuron axons, and why is there such a restricted pattern of RegIII β expression?

Axons regulate Schwann cells during development of the peripheral nervous system (29–33). The axon signals for regulating Schwann cell differentiation are known to include both proteins encoded by the neuregulin gene (NRG) signaling through the erbB family of receptors, adhesion molecules such as L1 and N-cadherin and neurotrophic factors such as brain-derived neurotrophic factor (BDNF) (34–37). It has been suggested that cell adhesion molecules on the axon surface as well as signals from the extracellular matrix and neurotrophins are also required for myelination to proceed efficiently and accurately. For example, BDNF supports postnatal facial motor neurones after axotomy and acts through p75 NTR to inhibit Schwann cell migration and promote myelination (35, 36, 38). It seems likely that RegIII β released from subsets of motor neuron axons is playing much the same role as BDNF in driving the process of myelination. As with NRG1 type III, it has been shown that RegIII β signals through the PI3-kinase pathway (8, 17), and this may represent a common pathway for axonally released RegIII β to influence Schwann cells. Why RegIII β expression is restricted to motor neurons concerned with the suckling and swallowing is unclear, but we suggest that the patterning of this

critical function in neonatal mice may be plastic and regulated by release of factors from target musculature concerned with suckling in an activity-dependent fashion (8).

Materials and Methods

All procedures complied with the United Kingdom Animals (Scientific Procedures) Act 1986.

Generation of the RegIII β -Deficient Mice by Gene Targeting. The generation of these mice has been described elsewhere (18). Complete knockout of the gene was confirmed in nervous tissue with immunocytochemistry by using antibodies generated against rat Reg2 protein, and absence of RegIII β mRNA was confirmed by using real-time PCR in brain, pancreas, and regenerating liver (16, 18).

Real-Time RT-qPCR Assay. Tissue preparation and RNA extraction were as described previously (39). RNA samples were treated with DNase I (Qiagen). Equal amounts (3 μ g) of total RNA were reversed transcribed by using random nonamers (Sigma) and SuperScript TM III RT (Invitrogen) for 1 h at 50°C in a total reaction volume of 20 μ l. cDNAs were immediately quantified by real-time PCR or kept at -20°C until further experiments. Real-time PCRs were performed with a DNA Engine (Bio-Rad) by using SYBR Green Jump Start RT-PCR master mix (Sigma) with each gene-specific primer (RegIII β forward, 5'-AAGAATATACCTCCGCACGC-3'; RegIII β reverse, 5'-CAGACAT-AGGGCAACTTCACC-3'; RegIII α forward, 5'-GAAGTGCCCTCTCCACGTACC-3'; RegIII α reverse, 5'-ACAAATGGTAATGTCCCATCG-3'; RegIII γ forward, 5'-GCTCCTATTGCTATGCTTGTAG-3'; RegIII γ reverse, 5'-CATGGAGGACAG-GAAGGAAGC-3'; β -actin forward, 5'-CAACGAGCGGTCCGATG-3'; β -actin reverse, 5'-GCCACAGGATTCCATACCA-3'). One microliter of cDNA was amplified in a three-step cycling program in a final reaction volume of 25 μ l. Control cDNA samples (obtained without transcriptase) were always included, as well as samples without any cDNA template. Reactions were performed in triplicate for five to six biological replicates, and threshold cycle values were normalized to β -actin gene expression. The specificity of the products was determined by melting curve analysis. The ratio of the relative expression of target genes to β -actin was calculated by using the $2^{-\Delta CT}$ formula.

Facial Nerve Section. Pups were cooled on wet ice (4°C), and the extent of anesthesia was determined by assessing reflex responses to tail pinch. Adult mice were anesthetized with Fluothane. The right facial nerve was transected at the stylomastoid foramen. One to 10 days later, mice were deeply anesthetized with Euthatal i.p. and transcardially perfused with 4% paraformaldehyde (PFA) in 0.1 M sodium phosphate buffer (PB) (pH 7.4) preceded by a brief wash with heparinized saline. After a 2 h postfix in PFA, tissue was moved to sucrose (30%) PB solution in preparation for freeze sectioning. In some pups, 5 μ l of CNTF (250 ng) applied to a piece of Gelfoam in 5 μ l saline, or saline alone, was applied to the cut end of the facial nerve at the time of section. Animals received only one treatment: either CNTF or saline.

Detection of β -Galactosidase Staining by X-Gal Staining. PFA-fixed tissue was washed in PBT (PBS/0.1% Tween-20) several times. After a rinse in X-Gal staining solution [5 mM K₃Fe(CN)₆, 5 mM K₄Fe(CN)₆, 1 mM MgCl₂, 0.01% sodium deoxycholate, 0.02% Triton plus X-Gal (4-chloro,5-bromo,3-indolyl-

β -galactosidase) at 1 mg/ml, suspended in PBS (pH 7.2)], tissue was transferred to fresh staining solution. Tissue was incubated at 37°C in the dark overnight. Sections were washed several times with PBT.

Cell Counting. Sections (40 μ m) were cut through the facial nucleus and all sections (the facial nucleus is \approx 700 μ m in length in the anterior-posterior direction) were mounted and stained with neutral red. All neuronal profiles in every section were counted and total counts corrected by using the Abercrombie correction (40).

Immunohistochemistry. Mice were deeply anesthetized with pentobarbitone (60 mg/kg, i.p.) and transcardially perfused with 4% PFA in 0.1 M sodium phosphate buffer (PB) (pH 7.4) preceded by a brief wash with heparinized saline. Tissue was dissected and postfixed for 2 h at 4°C and then transferred to 30% (wt/vol) sucrose in 0.1 M PB containing 0.02% (wt/vol) sodium azide. 40 μ m free-floating tissue sections were processed as previously described but for detection of RegIII β (9, 39). Hypoglossal innervation of tongue musculature in wild-type and KO mice was assessed with PGP, a marker for nerve fibres, and α -bungarotoxin, a marker for muscle end plates.

Electron Microscopy. Mouse pups or adult mice (P21) were deeply anesthetized with Nembutal and perfused with 10 ml heparinized saline followed by 20 ml 4% PFA plus 0.5% glutaraldehyde. Tissue was fixed overnight before dissection of the caudal tongue and attached hypoglossal nerves and the oculomotor nerve still attached to the optic nerve and eyecup. After being osmicated (30 min in 1% OsO₄ in 0.1 M PB), the sections were stained for 15 min in 0.1% uranyl acetate in sodium acetate buffer at 4°C, dehydrated in ethanol, cleared in propylene oxide, and embedded in Araldite. Semithin sections were cut with glass knives and stained with toluidine blue adjacent to thin sections cut with a diamond knife on an Ultracut ultramicrotome (Reichert). The sections were collected on mesh grids coated with a thin Formavar film, counterstained with lead citrate, and viewed in a JEOL 1010 electron microscope. Counts were made from four photographs of the lateral and medial nerves for each animal taken at \times 4,000 magnification. Total counts of myelinated fibers at P21 were made from photomicrographs of the lateral and medial hypoglossal nerves at \times 1,000 magnification.

Measurement of Milk/Colostrum Intake. The procedure described by Fujita and colleagues (41, 42) was followed. Briefly, pups were separated from the dam for 2 h and kept in a warm, dark chamber. The pups were then weighed and placed back with the mothers for 1 h before reweighing. Thirty to 40 pups of each genotype were used for the measurement of milk intake.

Statistical Analysis. The data are expressed as means \pm SEM. The data were analyzed by general linear model univariate or multivariate test, as appropriate, followed by Bonferroni post hoc tests or Student's *t* test, as appropriate. For all statistical analysis, statistical significance was set at *P* < 0.05.

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